

SEQUENCE LISTING

| <110> Europäisches Laboratorium für Molekularbiologie | | | | | | | | | | | | | | |
|---|---|--|--|--|--|--|--|--|--|--|--|--|--|--|
| 120% Protein with cell proliferation and cell division modulating activity and DNA encoding such protein | | | | | | | | | | | | | | |
| 1307 19595PWO | | | | | | | | | | | | | | |
| +146> PCT/EP00/00877 +141+ 2000-02-03 | | | | | | | | | | | | | | |
| 1150 - EP 39102172.6 1151: 1993-02-03 | | | | | | | | | | | | | | |
| 160:- 4 | | | | | | | | | | | | | | |
| -:170:- PatentIn Ver. 2.1 | | | | | | | | | | | | | | |
| +:210:- 1 -2:1: | | | | | | | | | | | | | | |
| +:22C:+ +:221:+ CDS +:222:- (214)(1113) | | | | | | | | | | | | | | |
| <:400 + 1 ttaaacagga cttgcagctc cagtgtaggt tttttcagaa getecgcecc aatgctgtat 60 | | | | | | | | | | | | | | |
| ttttttatta ttocaggagg ctataaagag agcagacaaa ggaagtaggc ggagttootg 12 | 0 | | | | | | | | | | | | | |
| tituatogoda titiggodagi ggiggotaag ogodigaagg iggoligotio olitigotoag 18 | 0 | | | | | | | | | | | | | |
| atcaacocto gggeoggtgt coccetttot ana ang agg cac atg cag agt gta 23 Met Arg His Met Gln Ser Val 1 \$ | 4 | | | | | | | | | | | | | |
| and tgg god ago too att tgt ggd ago ggg gtg aag dag gtd att ggd 28 | | | | | | | | | | | | | | |
| Thr Arg Ala Ser Ser Ile Cys Gly Ser Gly Val Lys Gln Val Ile Gly 10 15 20 | | | | | | | | | | | | | | |
| aag ggg cat ccg cac gcc cgg gtt gtt gga gcg cgc aag gcg caa atc 33 Lys Gly His Pro His Ala Arg Val Val Gly Ala Arg Lys Ala Gln Ile 25 30 35 | 0 | | | | | | | | | | | | | |
| set gag aga gag gag ttg tea gte aaa see aaa atg gtg ega aat ace 37 | 8 | | | | | | | | | | | | | |

| Pro 40 | Glu | Arg | Glu | Glu | Leu 45 | Ser | Val | Lys | Pro | Lys 50 | Met | Val | Arg | Asn | Thr 55 | |
|-----------|-----|-----|-----|-----|-----------|-----|-----|-----|-------------------|------------------|-----|-----|-----|-----|-----------|-------------|
| | | | | | | | | | cag Gir. 65 | | | | | | | 426 |
| | | | | | | | | | tot Ser | | | | | | | 474 |
| | | _ | _ | | | | - | _ | gtt Val | | - | | | | | 522 |
| | | | | | | | | | aca Thr | | | | | | | 570 |
| | | | | | | | | | gaa Glu | | | | | | | 618 |
| | | | | | | | | | gad Asp 145 | | | | | | | 66₽ |
| | | | _ | _ | | | - | | ttc Phe | | | | | | | 714 |
| | | | | | | | | | gat Asp | | | | | | | 76 <u>:</u> |
| | | | | | | | | | age Ang | | | | | | | 810 |
| | | | | | | | | | gac Asp | | | | | | | £5° |
| | | | | | _ | | | | ctt Leu 225 | | | | | | | 90 ń |
| tgt | gac | tct | ggt | ggg | gtc | tcc | cac | aac | aac | tst | tcc | tst | cca | gaa | caa | 954 |

| Cys Asp Ser Gly Gly Val Ser His Asn Asn Ser Ser Ser Pro Glu Gln 235 240 245 | |
|---|------|
| gag att tit dad tad add aat agg gag tigg tod dag gag oft ofd atg Glu Ile Phe His Tyr Thr Asn Arg Glu Trp Ser Gln Glu Leu Leu Met 250 255 260 | 1002 |
| tig occ bot gag otg tig otg gat occ gag tgt act cat gac tia cac Leu Pro Pro Glu Leu Leu Asp Pro Glu Cys Thr His Asp Leu His 265 270 275 | 1050 |
| att oto dag gag oda tig git gga tia gag oda gat ggg adg gog otg Ile Leu Gln Glu Pro Leu Val Gly Leu Glu Pro Asp Gly Thr Ala Leu 280 285 290 295 | 1098 |
| gaa tgg cac sac ott tagtagooga ttgtotooto ogagotttta ttottotota Glu Trp His His Leu 300 | 1153 |
| stoasaaget eageaettat teteteetee taaggaettig teaatigttsa gaettaattig | 1213 |
| aaatgggaga agtgaatatt उठ्युव्प्युप्यानु tagagtjijja नानां गावटाट agaga स्तनान्तु । | 1277 |
| ttttgagtot gtataaacog ttgstttgta aataaatata taaatgttot otgtgstggt | 1333 |
| cactaataaa gatcaggtaa aatcacttto aggtgtaatt taatagtatg tatgtagagt | 1393 |
| ottitaatica geteteeace aaatagtaae tigicateae igaacettig oftaactaca | 1453 |
| ettttattat tetgeacaca aatattetga agateagace gttetgtttt eagatgggtt | 1513 |
| gaaaatatta aactcaacag aatteetgtg gtgtaatgta aatgcaaaga tegattagac | 1573 |
| ^L ā | 1575 |

(213> Xenopus sp.

<400> 2

Met Arg His Met Gln Ser Val Thr Arg Ala Ser Ser Ile Cys Gly Ser 1 5 10 15

Gly Val Lys Gln Val Ile Gly Lys Gly His Pro His Ala Arg Val Val
20 25 30

Gly Ala Arg Lys Ala Gln Ile Pro Glu Arg Glu Glu Leu Ser Val Lys 3.5 4.0 Pro Lys Met Val Arg Asn Thr His Lew Asn Lew Gln Pro Gln Glu Arg 5.5 60 Gln Ala Phe Tyr Arg Leu Leu Glu Asn Glu Gln Ile Gln Glu Phe Leu 70 75 Ser Met Asp Ser Cys Leu Arg Ile Ser Asp Lys Tyr Leu Ile Ala Met 85 90 95 Val Leu Ala Tyr Phe Lys Arg Ala Ala Gly Leu Tyr Thr Ser Glu Tyr 105 100 Thr Thr Met Asn Phe Phe Val Ala Leu Tyr Leu Ala Asn Asp Met Glu 120 125 115 Glu Asp Glu Glu Asp Tyr Lys Tyr Glu Ile Phe Pro Trp Ala Leu Gly 130 135 Asp Ser Trp Arg Glu Leu bne Pro Glo The Leu Arg Leu Arg Asp Asp 155 145 150 Phe Trp Ala Lys Met Asn Tyr Arg Ala Val Val Ser Arg Arg Cys Cys 165 170 Asp Glu Val Met Ser Lys Asp Pro Thr His Trp Ala Trp Leu Arg Asp 180 185 Arg Pro Met His His Ser Gly Ala Met Arg Gly Tyr Leu Arg Asn Glu 195 200 205 Asp Asp Phe Pro Arg Gly Pro Gly Lea Thr Pro Ala Ser Cys Thr 215 220 In: The His Eys Ala Gly Val Cys Asp Ser Gly Gly Val Ser His Asn 225 230 235 Asn Ser Ser Pro Glu Gln Glu Ile Phe His Tyr Thr Asn Arg Glu 245 250 Trp Ser Gln Glu Leu Leu Met Leu Pro Pro Glu Leu Leu Asp Pro 265 Glu Cys Thr His Asp Leu His Ile Leu Gln Glu Pro Leu Val Gly Leu 275 280 285

| Glu | Pro | Asp | Gly | Thr | Ala | Leu | Glu | Trp | His | His | Leu |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| | 290 | | | | | 295 | | | | | 300 |

| (210)(3) | | | | | | | | | | | | | | |
|--|--|--|--|--|--|--|--|--|--|--|--|--|--|--|
| -(211): 1357 | | | | | | | | | | | | | | |
| 1212 - DNA | | | | | | | | | | | | | | |
| - 213 - Kenopus sp. | | | | | | | | | | | | | | |
| | | | | | | | | | | | | | | |
| (220) | | | | | | | | | | | | | | |
| 42212 ODS | | | | | | | | | | | | | | |
| -7222> (163)(1056) | | | | | | | | | | | | | | |
| | | | | | | | | | | | | | | |
| H(400)H 3 | | | | | | | | | | | | | | |
| gotgtatttt actttottto aggaggotat aaagacaaca gacaggggag gtaggcagag 6 | | | | | | | | | | | | | | |
| | | | | | | | | | | | | | | |
| ttootgttoa toaccattot ttggoogttg gtggotaggo gootgaaggt ggotgotado 120 | | | | | | | | | | | | | | |
| | | | | | | | | | | | | | | |
| tttgotbaga hoaadootog yttogttgto odeotttota oa aty agy dat atg = 174 = | | | | | | | | | | | | | | |
| Met Arg His Met | | | | | | | | | | | | | | |
| | | | | | | | | | | | | | | |
| cad agt goa ace egg goe ace tta gtt tgt gge age ggg gta aag cag = 222 | | | | | | | | | | | | | | |
| cag agt goa acc ogg god acc tta gtt tgt ggd agd ggg gta aag dag - 222 - Gln Ser Ala Thr Arg Ala Thr Leu Val Cys Gly Ser Gly Val Lys Gln | | | | | | | | | | | | | | |
| 5 10 15 20 | | | | | | | | | | | | | | |
| | | | | | | | | | | | | | | |
| Atc att god aag gga cat deg aat acc ogg gtt tit gga geg ogd aag - 270 | | | | | | | | | | | | | | |
| The The Ala Lys Gly His Pro Asn Thr Arg Val Phe Gly Ala Arg Lys | | | | | | | | | | | | | | |
| 25 30 35 | | | | | | | | | | | | | | |
| | | | | | | | | | | | | | | |
| gog aaa ato oot gag aga gag gtg ota goa goo aaa ooc aag ato acg - 318 | | | | | | | | | | | | | | |
| Ala Lys Ile Pro Glu Arg Glu Val Leu Ala Ala Lys Pro Lys Ile Thr | | | | | | | | | | | | | | |
| 40 45 50 | | | | | | | | | | | | | | |
| | | | | | | | | | | | | | | |
| ogo att ada dat otd dat otd bud pod hag gud ogo bag god tit tad 366 | | | | | | | | | | | | | | |
| Act I'm Thr His Leu Asn Leu Gln Pro Gln Glu Arg Gln Ala Phe Tyr | | | | | | | | | | | | | | |
| စ်ပ | | | | | | | | | | | | | | |
| | | | | | | | | | | | | | | |
| agg ofc one gas ast gag ofg att dag gas tit oft tot atg gad tid 4.4 | | | | | | | | | | | | | | |
| Arg Leu Leu Glu Asn Glu Leu Ile Gln Glu Phe Leu Ser Met Asp Ser 70 75 80 | | | | | | | | | | | | | | |
| 7.5 75 80 | | | | | | | | | | | | | | |
| tgt ota aag att toa gas aag tat oto ata goa atg gtt ota goa tat = 462 | | | | | | | | | | | | | | |
| Cys Leu Lys Ile Ser Asp Lys Tyr Leu Ile Ala Met Val Leu Ala Tyr | | | | | | | | | | | | | | |
| 85 90 95 100 | | | | | | | | | | | | | | |
| | | | | | | | | | | | | | | |
| ttt aag ogg gog ggo oto tac acc ago gag tac aca acc atg aat tto - 510 | | | | | | | | | | | | | | |
| | | | | | | | | | | | | | | |

| Ehe | Lys | Arg | Ala | Gly 105 | Leu | Tyr | Thr | Ser | Glu 110 | Tyr | Thr | Thr | Met | Asn 115 | Phe | |
|-----|-----|-----|-----|------------|-----|------|-------|-------------------|------------|------|-------|------|------|------------|-----|------|
| | | - | | | _ | _ | | gae Asp 125 | - | | _ | - | - | | | 558 |
| | | | - | | | | | gca Ala | | | | | | _ | | 60-6 |
| | | | | | | - | | ogg Arg | _ | _ | | | - | | - | 654 |
| | | | | | | | | aga Arg | | | | | | | | 702 |
| | | | | | | | | ats Lea | | | | | | | | 750 |
| _ | | _ | _ | - | | | | aga Arg 205 | | | | _ | | | | 798 |
| | | | | | | | | agc Ser | | | | | | | | 84ნ |
| _ | - | - | - | | | | | too Ser | | • | | | | | - | 894 |
| | | | | | | | | agg Arg | | | | | | | | 942 |
| | | | | | | | | gat Asp | | | | | | | | 990 |
| | | | - | - | - | _ | - | gga Gly 285 | | | | | | - | | 1038 |
| ttg | gaa | tgg | cac | cac | ctt | tage | cacca | atg t | cato | etet | gt go | cttt | catt | 2 | | 1086 |

Leu Glu Trp His His Leu 295

ttotoraato cacqaqetea agaageaett aaeeteteet aageaettge chatgteeet 1146 attoagasta atgaattaaa tgggagaggt gactattgee ataaagggaa ggatgeeaet 1206 tagagtggag aataataett geeaaaaatg gtgtttgggt etgtttaaae tgttgetatt 1266 teagttgeet tgtaaataaa taagtataaa aatgtatget etgtgeeggt tgetaataaa 1326 aaaaaaatet ggtateaaaa aaaaaaaaa a 1357

<2105 4

·211: 298

12121 PRT

3213 Menopus sp.

· 400:- 4

Met Ard His Met Gln Ser Ala Thr Arg Ala Thr Leu Val Cys Gly Ser

Gly Val Lys Gln Ile Ile Ala Lys Gly His Pro Asn Thr Arg Val Phe
20 25 30

Gly Ala Arg Lys Ala Lys Ile Pro Glu Arg Glu Val Leu Ala Ala Lys 35 40 45

Fro Lys Ile Thr Arg Ile Thr His Leu Asn Leu Gln Pro Gln Glu Arg 50 60

Gln Ala Phe Tyr Arg Leu Leu Glu Asn Glu Leu Ile Gln Glu Phe Leu 65 70 75 80

Ser Met Asp Ser Cys Leu Lys Ile Ser Asp Lys Tyr Leu Ile Ala Met 85 90 95

Val Leu Ala Tyr Fne Lys Arg Ald Gly Leb Tyr Thr Ser Glu Tyr Thr 100 105 110

Thr Met Asn Phe Phe Val Ala Leu Tyr Leu Ala Asn Asp Met Glu Glu
115 120 125

Asp Gld Glu Asp Tyr Lys Tyr Glu Ile Phe Pro Trp Ala Leu Gly Asp 130 140

Ser Trp Arg Glu Phe Phe Pro Gln Phe Leu Arg Leu Arg Asp Asp Phe

- Trp Ala Lys Met Asn Tyr Arg Ala Val Val Ser Arg Arg Cys Cys Asp 165 170 175
- Glu Val Met Ala Lys Asp Pro Thr His Trp Ala Trp Leu Arg Asp Arg 180 185 190
- Pro Ile His His.Ser Gly Ala Leu Arg Gly Tyr Leu Arg Asn Glu Asp 195 200 205
- Asp Phe Phe Pro Arg Gly Pro Gly Leu Thr Pro Ala Ser Cys Ala Leu 210 215 220
- Cys His Lys Ala Ser Val Cys Asp Ser Gly Gly Val Ser His Asp Asn 225 230 235 240
- Ser Ser Pro Glu Gln Glu Ile Phe His Tyr Thr Asn Arg Glu Trp Ser 245 250 255
- Alt Alt Leu Leu Fle Leu Pro Pro Glu Leu Leu Leu Asp Pro Glu Ser 260 Leu
- Thr Tyr Asp Ile His Ile Phe Gln Glu Pro Leu Val Gly Leu Glu Pro 275 280 285
- Asp Gly Ala Ala Leu Glu Trp His His Leu 290 295

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SEQ ID NO. 1

ls 26 cDNA

TTA AAC AGG ACT TGC AGC TCC AGT GTA GGT TTT TTC AGA AGC TCC GCC CCA ATG CTG TAT TTT TTT ATT ATT CCA GGA GGC TAT ANA GAG AGC AGA CAA AGG AAG TAG GCG GAG TTC CTG 61 121 TTT ATC GCC ATT TGG CCA GTG GTG GCT AAG CGC CTG AAG GTG GCT GCT TCC TTT GCT CAG 181 ATC AAC CCT CGG GCC GGT GTC CCC CTT TCT ACA ATG AGG CAC ATG CAG AGT GTA ACC CGG H M 241 GCC AGC TCC ATT TGT GGC AGC GGG GTG AAG CAG GTC ATT GGC AAG GGG CAT CCG CAC GCC ĸ G CGG GTT GTT GGA GCG CGC AAG GCG CAA ATC CCT GAG AGA GAG GAG TTG TCA GTC AAA CCC 361 AAA ATG GTG CGA AAT ACC CAT CTC AAT CTA CAG CCC CAG GAG CGC CAG GCC TTC TAC AGG 421 CTC CTA GAA AAT GAG CAG ATT CAG GAA TTC CTT TCT ATG GAC TCC TGT CTA AGG ATT TCC 0 Ξ М 481 GAC AAG TAT CTC ATA GCA ATG GTT CTA GCA TAT TTT AAG CGG GCA GCG GGC CTC TAC ACC λ 541 AGC GAG TAC ACA ACC ATG AAT TTC TTT GTT GCC CTG TAT CTG GCT AAT GAC ATG GAA E 129 601 GAT GAA GAA GAC TAT AAA TAT GAA ATC TTC CCC TGG GCA CTA GGA GAC TCG TGG CGT GAG R 661 CTT TTC CCA CAA TTT TTG CGT CTC CGG GAC GAC TTC TGG GCT AAA ATG AAC TAC CGA GCA 169 721 GTT GTT AGT CGA AGG TGC TGT GAT GAG GTA ATG TCC AAA GAT CCC ACT CAT TGG GCC TGG 189 λ 781 CTG AGA GAT CGC CCC ATG CAT CAC AGC GGG GCC ATG CGT GGT TAC CTT AGA AAC GAG GAC н Н R R N E 209 841 GAC TIT TIC CCC CGG GGT CCA GGC CIT ACA CCA GCC AGC TGT ACA CIT TGC CAT AAA GCA R G 229 901 GGT GTC TGT GAC TCT GGT GGG GTC TCC CAC AAC AAC TCT TCC TCT CCA GAA CAA GAG ATT Ħ 961 TTT CAC TAC ACC AAT AGG GAG TGG TCC CAG GAG CTT CTC ATG TTG CCC CCT GAG CTG TTG 269 0 Ε 1021 CTG GAT CCC GAG TGT ACT CAT GAC TTA CAC ATT CTC CAG GAG CCA TTG GTT GGA TTA GAG D E D H I E P T. v G L E 1081 CCA GAT GGG ACG GCG CTG GAA TGG CAC CTT TAG TAG CCG ATT GTC TCC TCC GAG CTT 300 1141 TTA TTC TTC TCT ACT CAC AAG CTC AGC ACT TAT TCT CTC CTC CTA AGG ACT TGT CAA TGT 1201 TCA GAC TTA ATT GAA ATG GGA GAA GTG AAT ATT CCG ACG GAT GTA GAG CGG GAA TAT GTG 1261 CCC AGA GAA AGT GTT TTG AGT CTG TAT AAA CCG TTG CTT TGT AAA TAA ATA TAT AAA TGT 1321 TCT CTG TGC TGG TCA CTA ATA AAG ATC AGG TAA AAT CAC TTT CAG GTG TAA TTT AAT AGT 1381 ATG TAT GTA GAG TOT TTA ATT CAG CTC TCC ACC AAA TAG TAA CTT GTC ATC ACT GAA CCT 1441 TTG CTT AAC TAC ACT TIT ATT ATT CTG CAC ACA AAT ATT CTG AAG ATC AGA CCG TTC TGT 1501 TTT CAG ATG GGT TGA AAA TAT TAA ACT CAA CAG AAT TCC TGT GGT GTA ATG TAA ATG CAA 1561 AGA TCG ATT AGA CTA

SEQ ID NO. 2

ls 27 cDNA

GCT GTA TTT TAC TTT CTT TCA GGA GGC TAT AAA GAC AAC AGA CAG GGG AGG TAG GCA GAG TTC CTG TTC ATC ACC ATT CTT TGG CCG TTG GTG GCT AGG CGC CTG AAG GTG GCT ACC 121 TTT GCT CAG ATC AAC CCT CGG TTC GTT GTC CCC CTT TCT ACA ATG AGG CAT ATG CAG AGT R H 181 GCA ACC CGG GCC ACC TTA GTT TGT GGC AGC GGG GTA AAG CAG ATC ATT GCC AAG GGA CAT 26 241 CCG AAT ACC CGG GTT TTT GGA GCG CGC AAG GCG AAA ATC CCT GAG AGA GAG GTG CTA GCA Ε 301 GCC AAA CCC AAG ATC ACG CGC ATT ACA CAT CTC AAT CTA CAA CCC CAG GAG CGC CAG GCC H 0 361 TTT TAC AGG CTC CTA GAA AAT GAG CTG ATT CAG GAA TTT CTT TCT ATG GAC TCC TGT CTA N 421 ANG ATT TON GAC ANG THT CTC ATA GON ATG GTT CTN GON THT TIT ANG CGG GGG GGC CTC G 106 481 TAC ACC AGC GAG TAC ACA ACC ATG AAT TTC TTT GTT GCT CTG TAT CTG GCT AAT GAC ATG 126 541 GAG GAA GAT GAA GAA TAT AAA TAT GAA ATC TTC CCC TGG GCA CTA GGA GAT TCA TGG 501 CGT GAG TTT TTC CCA CAA TTT TTA CGT CTC CGG GAC GAC TTC TGG GCT ANA ATG AAC TAC 166 N 661 CGA GCA GTT GTT AGC CGA AGA TGT TGT GAT GAG GTA ATG GCG AAA GAT CCC ACT CAT TGG н 186 721 GCC TGG CTC AGA GAT CGT CCT ATT CAT CAT AGT GGG GCC CTG CGT GGT TAC CTC AGA AAT H 206 781 GAG GAT GAC TIT TTC CCT CGG GGT CCA GGC CTT ACA CCA GCC AGC TGT GCA CTT TGC CAT 841 AAA GCA AGT GTC TGT GAC TCT GGT GGG GTG TCC CAT GAC AAC TCT TCT CCA GAA CAA GAG E 0 901 ATT TTT CAC TAC ACC AAT AGG GAG TGG TCC CAG GAA CTT CTC ATC TTG CCA CCT GAA CTG 266 E 961 TTA TTG GAT CCG GAG TCT ACT TAT GAC ATC CAC ATT TTC CAG GAA CCG TTG GTT GGA TTA G 1921 GAG CCA GAT GGG GCA GCC TTG GAA TGG CAC CAC CTT TAG CAC CAT GTC ATC TCT GTG CTT н H 298 1081 TCA TTC TCT TAT CCA CGA GCT CAA GAA GCA CTT AAC CTC TCC TAA GCA CTT GCC CAT 1141 GTC CCT ATT CAG ACT AAT GAA TTA AAT GGG AGA GGT GAC TAT TGC CAT AAA GGG AAG GAT 1201 GCC ACT TAG AGT GGA GAA TAA TAC TTG CCA AAA ATG GTG TTT GGG TCT GTT TAA ACT GTT 1261 GCT ATT TCA GTT GCC TTG TAA ATA AAT AAG TAT AAA AAT GTA TGC TCT GTG CCG GTT GCT 1321 ANT ANA ANA ARA ATC TGG TAT CAN ANA ANA ANA ANA A



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| λ | s | s | I | C | G | s | G | v | ĸ | Q | v | r | G | ĸ | G | н | P | н | λ | 29 |
| R | ٧ | v | G | λ | R | ĸ | λ | Q | I | P | E | R | Ξ | E | L | s | v | ĸ | P | 49 |
| к | М | v | R | N | Ŧ | н | L | N | L | Q | P | Q | Ξ | R | Q | A | F | Y | R | 69 |
| L | L | E | N | E | Q | I | Q | £ | F | L | s | M | Ď | s | С | L | Ř | I | s | 89 |
| D | ĸ | Y | L | I | A | M | v | L | A | Y | F | ĸ | R | A | A | G | L | Y | T | 109 |
| s | Ε | Y | Ţ | Ť | M | N | F | F | v | A | L | Y | L | λ | N | D | м | E | E | 129 |
| D | Ε | Ξ | ā | Y | ĸ | Y | Ε | I | F | P | W | A | L | G | ۵ | s | W | R | Ξ | 149 |
| L | F | P | Q | F | L | R | L | R | α | D | F | w | λ | ĸ | М | N | Y | R | A | 169 |
| v | v | s | R | R | c | c | ۵ | E | v | M | s | к | D | P | Ŧ | H | W | λ | W | 189 |
| L | R | D | R | P | M | н | н | s | G | A | M | R | G | Y | L | R | N | E | a | 209 |
| Œ | F | F | P | R | G | P | G | L | Ŧ | P | A | s | С | T | L | C | H | ĸ | A | 229 |
| G | v | C | D | s | G | G | v | s | H | N | N | s | s | s | P | E | Q | E | I | 249 |
| F | н | Y | T | N | R | E | W | s | Q | E | L | L | M | L | P | P | E | L | L | 269 |
| L | D | ₽ | E | C | Ŧ | н | ם | L | H | I | L | Q | Σ | P | L | v | G | L | E | 289 |
| P | ۵ | G | T | λ | L | E | w | н | н | L | • | • | | | | | | | | 300 |

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M R н M s 6 R C G s G ĸ Q G 26 N G ĸ I 2 E R E v λ R ĸ L A 46 ĸ ₽ N Q Q E н 66 Y L E N Ε L I Q E M С 86 ĸ s ם v Y I ĸ ĸ R G I A L 106 Y L A М A Y т s L, v E Y M N F Α Y L A N D M 126 Ę ۵ Ε ĸ Ξ ₽ L G D s 146 R E F ם R D A N 166 ₽ Q L R L M R. v v P 186 A V s С C D Ε М λ ĸ D T H G 206 A W L I H H s G Y L R E D D F P R G P G T λ С C 226 ĸ E 246 s v v D N s s P E λ C ם s G G s н Q F H s Q Ε P Ε 266 L D D I H I Q 286 Ε s Y 298 P G E н н